



ARTIFICIAL INTELLIGENCE FOR NEW DRUG DESIGN

DEEP LEARNING FOR LIGAND-BASED DE
NOVO DESIGN IN LEAD OPTIMIZATION:
A REAL LIFE CASE STUDY



Iktos facts

Our company

Paris-based AI company founded late 2016

30 people

Specializing in AI applied to chemistry:

- Deep Generative models for *de novo* drug design
- Data-driven retrosynthesis

Concrete real-life experience of delivering value to drug discovery programmes: ~15 projects delivered or in progress

Business model: services, research collaborations, software

Our customers

 MERCK

 janssen
PHARMACEUTICAL COMPANIES
OF Johnson & Johnson

 GRÜNENTHAL

 SERVIER

 ORION

 ucb

 syngenta

 Galápagos

 Pierre Fabre

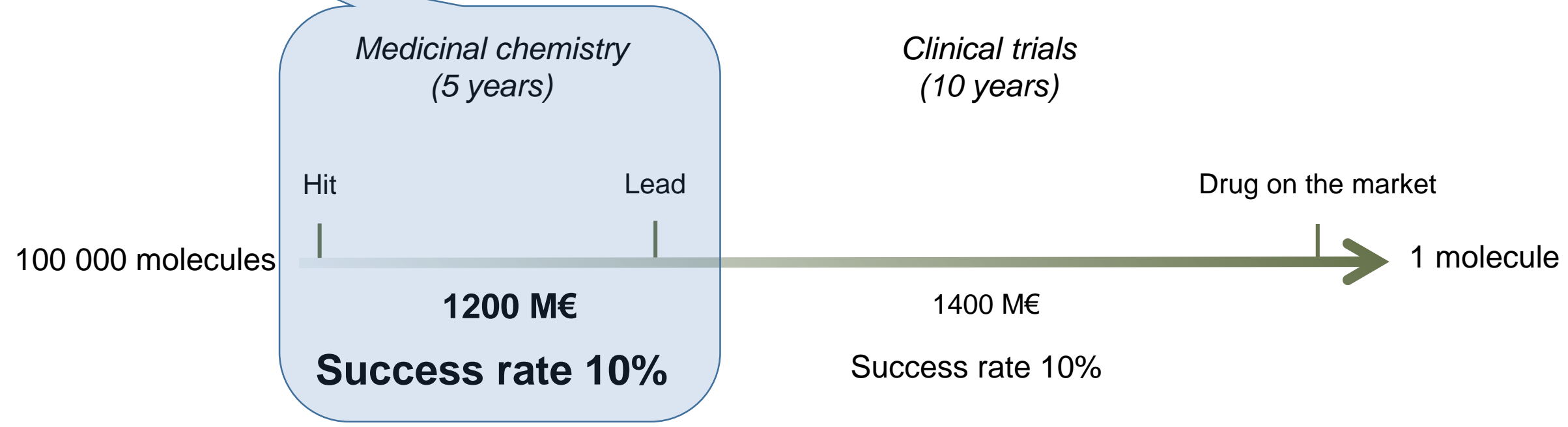
 DNDi
Drugs for Neglected Diseases initiative



MedChem: A difficult job!

life cycle: 10

Our focus



Less than 1% chance to find a new drug...

→ Our objective: use AI to identify the optimal molecule, faster, cheaper, with higher probability of success!

Solving the Rubik's cube:

- Simultaneous optimization on activity, potency, ADME, tox, selectivity...

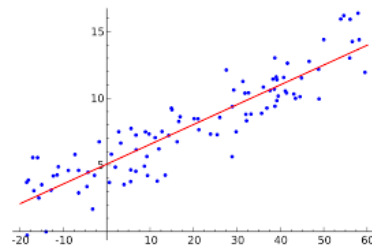


☹️ : Gain on one objective usually results in loss on the other ones

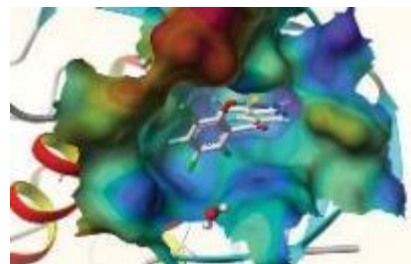
- The chemical space is huge (10^{60}). Does the solution even exist? Can we ever find it?

Predictive approaches:

- QSAR, data science

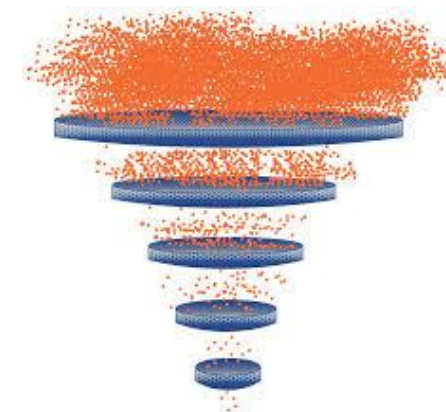


- Molecular modeling



Those technologies can only predict:
They tell you what not to do, but they do not tell you what to do!

Compound *de novo* design approaches:



Virtual screening: **Brute-force**

- ⊕ Limited by computational power and space
- ⊕ Only very small portions of the chemical space are explored (10^9 vs 10^{60})
- ⊕ Virtually Zero chance of finding “the” molecule

Evolutionary algorithms

- ⊕ Slow, limited diversity, compound feasibility issues

A filtering, rather than designing approach!



Artificial
intelligence:
changing the rules
of the game!

Deep Neuronal Net



Automatic colorization of black and white images



"girl in pink dress is jumping in air."



"black and white dog jumps over bar."



"young girl in pink shirt is swinging on swing."

Automatic image caption generation



Automatic game playing



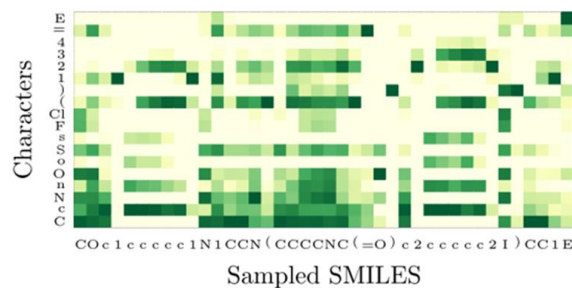
Automatic picture generation

Why not generate molecules instead of images of cats?

Generative

Sequence based

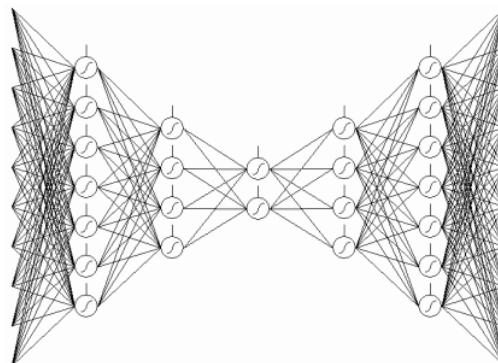
- Long Short Term Memory
- LSTM (Characters or Grammar)
- LSTM with Reinforcement learning



AstraZeneca 2017

Latent space

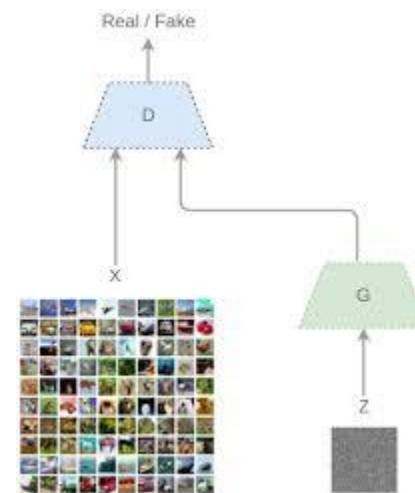
- Variational Auto-Encoder
- CVAE, GVAE, SD-VAE



Toronto
Harvard 2016

GAN

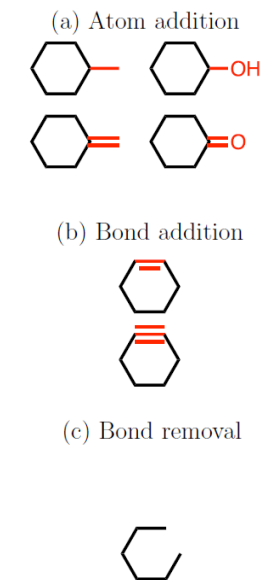
- Generative Adversarial Network
- ORGAN



Insilico Medicine 2017

DQN

- Deep Q-Networks



Google 2018

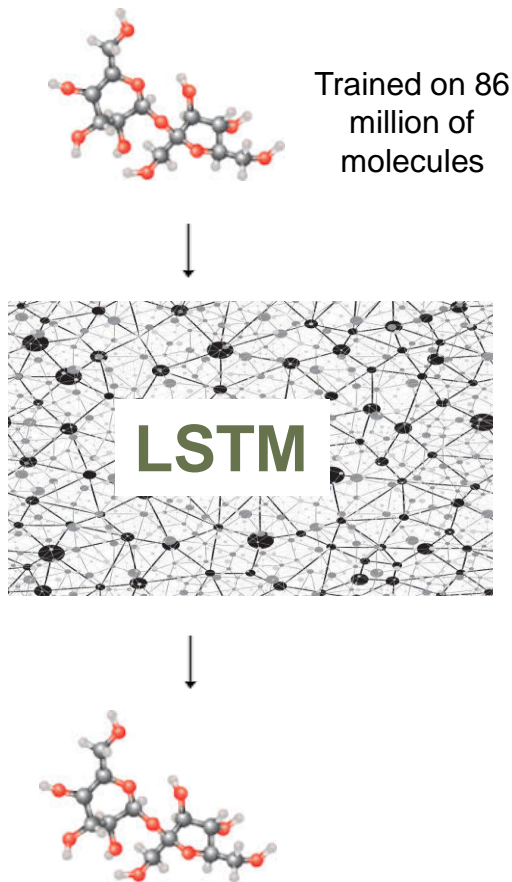
Recent review: Hongming Chen, Ola Engkvist, Yin Hai Wang, Marcus Olivecrona and Thomas Blaschke, *The rise of deep learning in drug discovery* Drug Discovery Today, 1-10, February 2018; Youjun Xu, Kangjie Lin, Shiwei Wang, Lei Wang, Chenjing Cai, Chen Song, Luhua Lai, Jianfeng Pei, *Deep learning for molecular generation*, FutureMed. Chem. January 2019; Daniel C. Elton, Zois Boukouvalas, Mark D. Fuge, Peter W. Chung, *Deep learning for molecular generation and optimization - a review of the state of the art*, arXiv March 2019

for MPO: LSTM

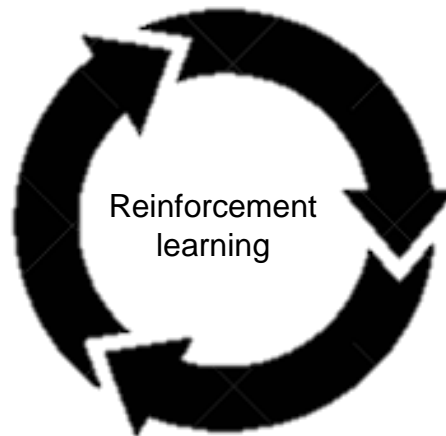
Generative model (AI)

Policy Gradient (AI)

- ✓ QSAR models
 - ✓ Docking score
 - ✓ Metrics, descriptors
 - ✓ Sub-structures
 - ...
- Traditional approaches

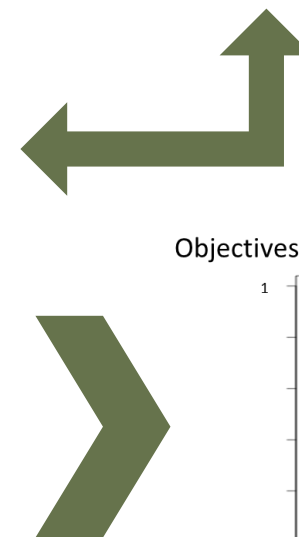


1) Molecules are generated

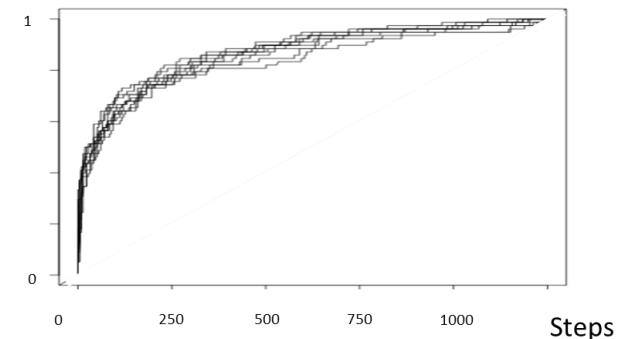


2) molecules are scored by the multi-objective fitness function

3) the weights of the model are adjusted to maximize the probability of generating molecules similar to those maximizing the global score using a policy gradient algorithm.

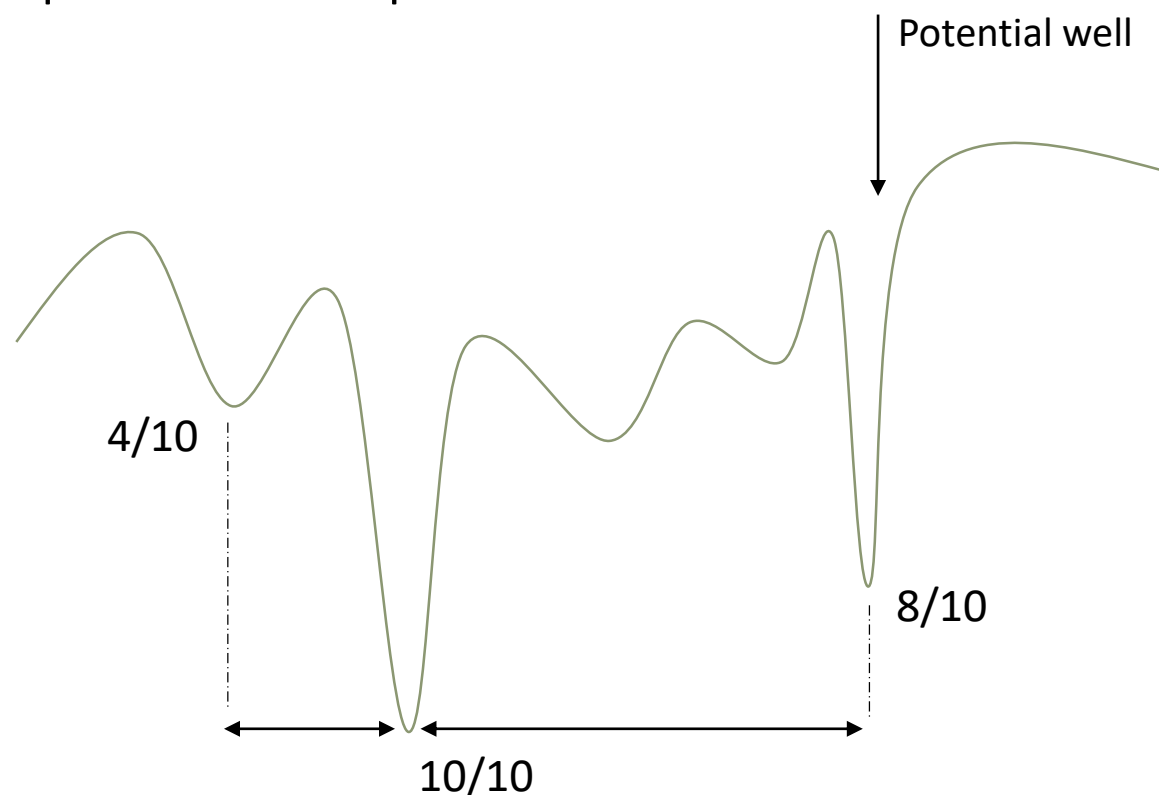


Objectives



Major MPO

10 parameters to optimize for instance



- Valid SMILES rate¹
- Synthetic access, compound “quality”
- QSAR Applicability Domain
- Diversity / Novelty²
- Global fitness function... or not..

A molecule which fixes 8 parameters over 10, doesn't mean it is closer to the solution than a molecule which fixes only 4 parameters!

1 – Mariya Popova, Olexander Isayev, Alexander Tropsha *Deep Reinforcement Learning for De-Novo Drug Design* arXiv Nov 2017

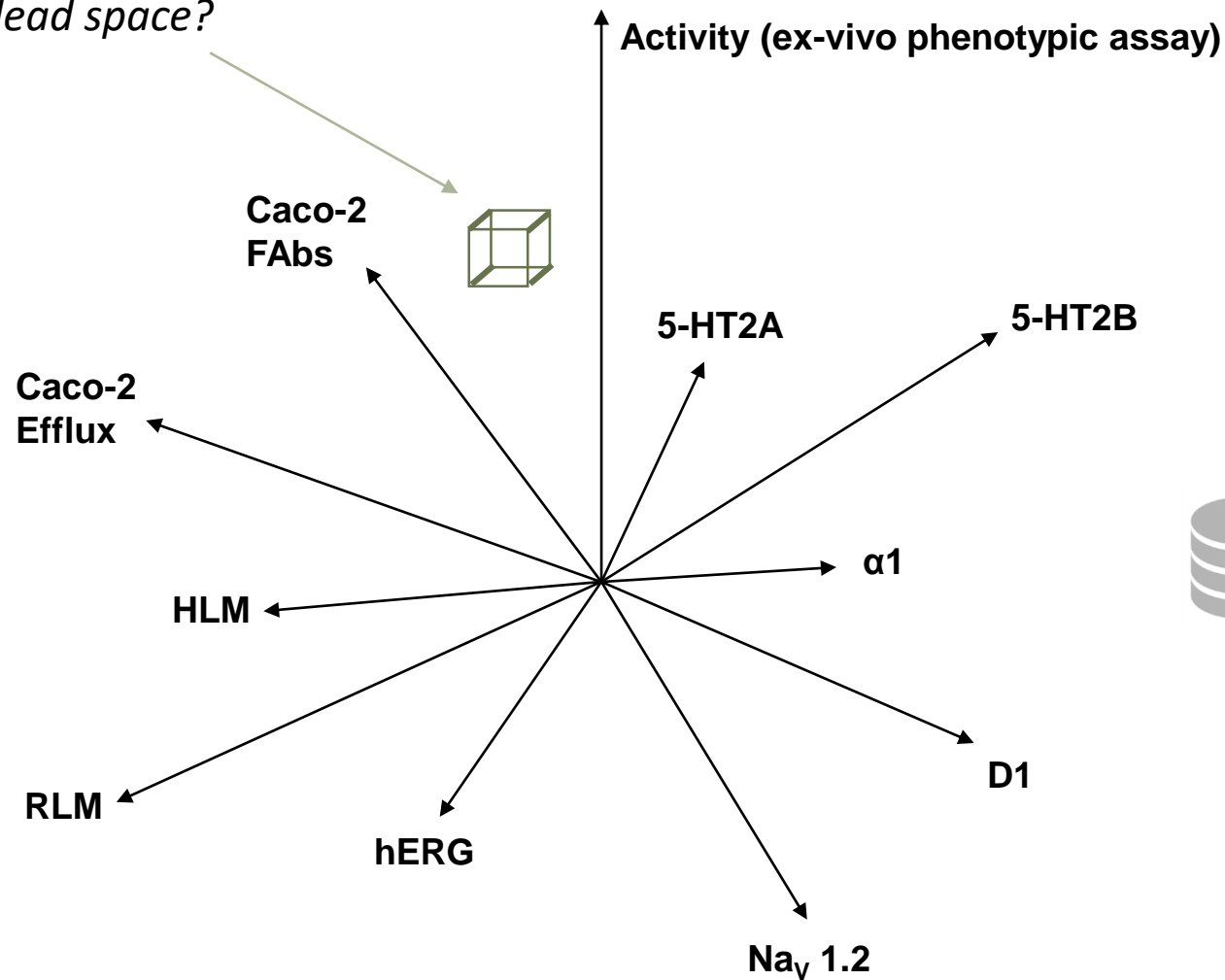
2 – Kristina Preuer, Philipp Renz, Thomas Unterthiner, Sepp Hochreiter, Günter Klambauer *Fréchet ChemblNet Distance: A metric for generative models for molecules* arXiv March 2018



Real life MPO Case study

PO Project cl

Where is the Optimal lead space?

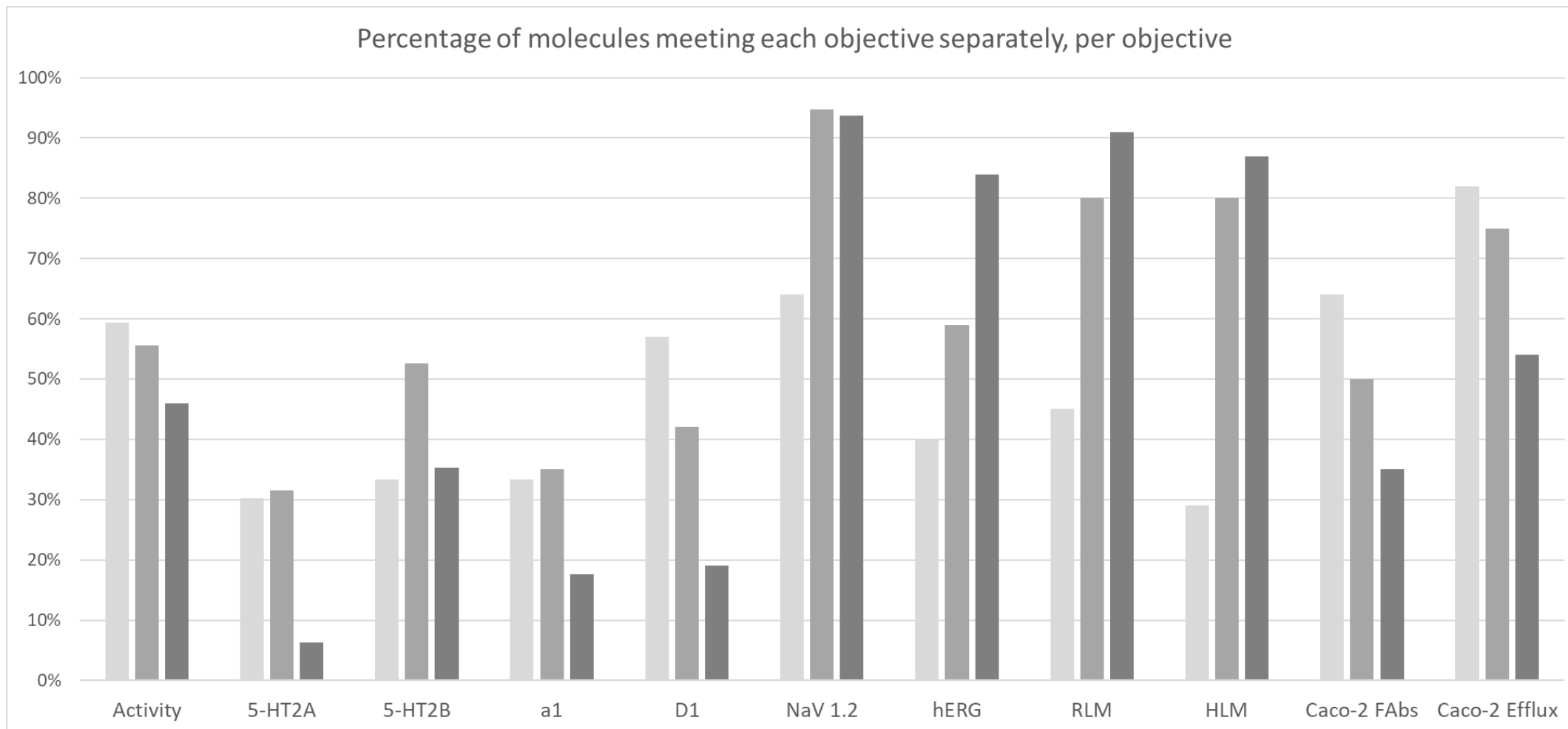


Project facts and figures

- Late-stage Lead Optimization Project
- Undisclosed target, complex, expensive ex-vivo phenotypic assay
- 11 objectives
- 880 molecules (251 measured on Activity)
- Several years of research

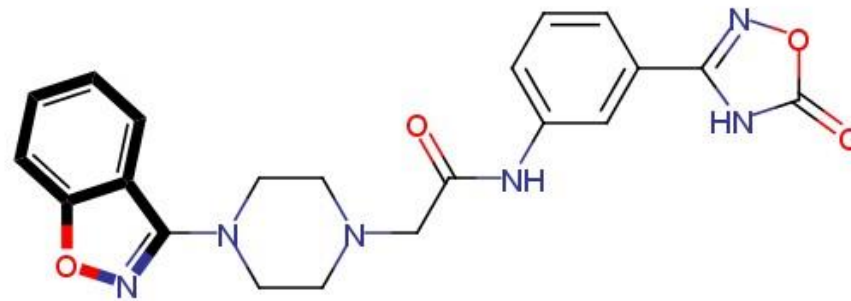
➔ *No molecule meeting simultaneously the 11 objectives of the blueprint...*

Dashboard also



Compound in t

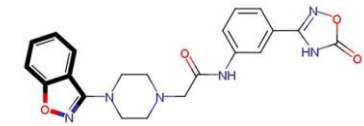
Activity	5-HT2A	5-HT2B	a1	D1	Na _v 1.2	hERG	RLM	HLM	Caco-2 FAbs	Caco-2 Efflux
194.0	20.0	18.0	1.0	4.0	0.0	19.0	82.85	63.35	88.99	26.2



Presence of a 1,2-benzoxazole moiety which appears in 61% of cases in the whole dataset and in 78% of the last 50 molecules

Compounds:

Activity	5-HT2A	5-HT2B	α 1	D1	Na _v 1.2	hERG	RLM	HLM	Caco-2 FAbs	Caco-2 Efflux
194.0	20.0	18.0	1.0	4.0	0.0	19.0	82.85	63.35	88.99	26.2
83.0	69.0	-25.0	45.0	6.0	13.0	6.4	69.04	31.93	97.6	1.96
46.0	46.0	69.0	14.0	14.0	-14.0	25.8	60.28	25.43	98.86	0.75
48.0	71.0	48.0	12.0	14.0	39.0	25.0	68.83	33.58	99.37	0.39
115.0	76.0	15.0	37.0	-3.0	-13.0	5.4	80.82	83.54	72.24	12.3
46.0	6.0	44.0	29.0	-11.0	20.0	12.4	93.11	78.36	73.8	34.1



mol 732

mol 663

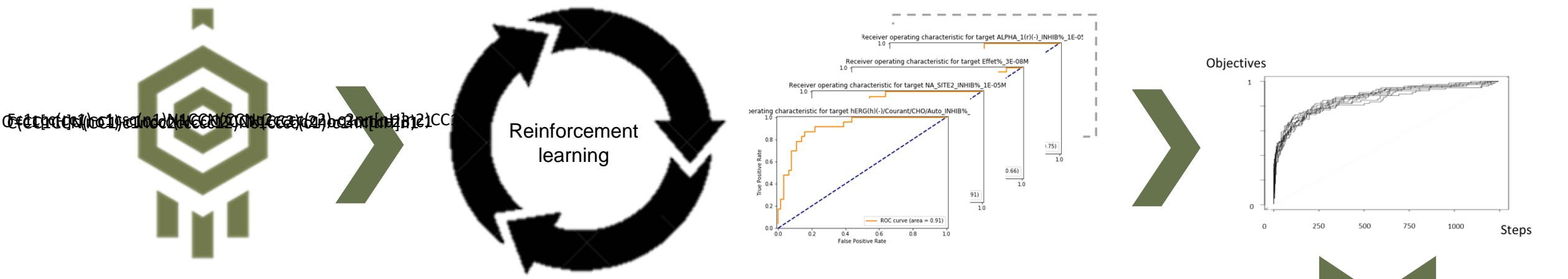
mol 559

mol 555

mol 550

mol 435

Virtual compound



Iktos molecules generator

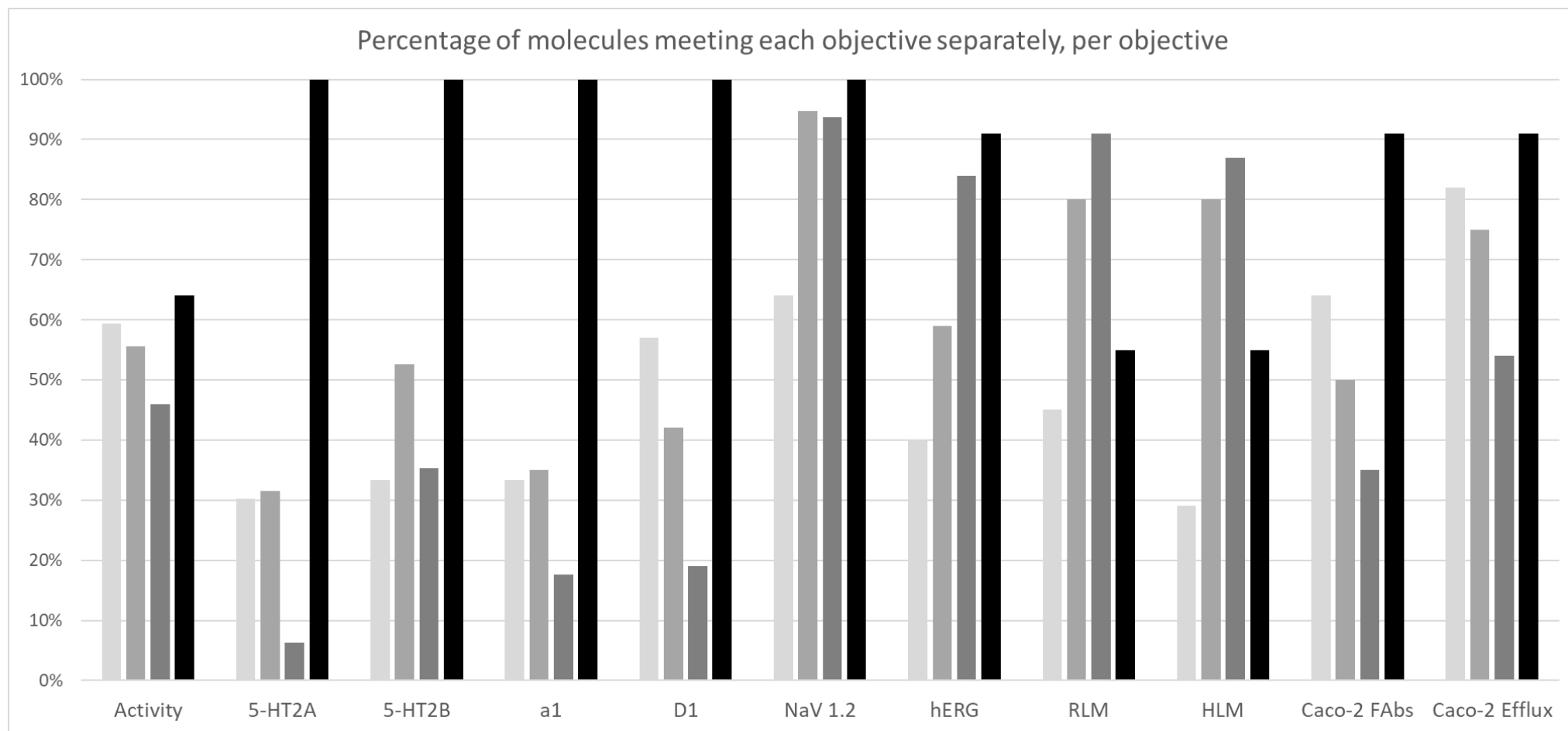
11 Independent QSAR models

Predicted Activity	Predicted 5-HT2A	Predicted 5-HT2B	Predicted α 1	Predicted D1	Predicted Na _v 1.2	Predicted hERG	Predicted RLM	Predicted HLM	Predicted Caco-2 FAbs	Predicted Caco-2 Efflux
Green	Green	Green	Green	Yellow	Red	Red	Red	Yellow	Yellow	Green
Green	Green	Green	Green	Yellow	Red	Green	Green	Yellow	Yellow	Green
Green	Green	Green	Green	Green	Green	Green	Green	Green	Green	Green

150 virtual “ideal” compounds were designed

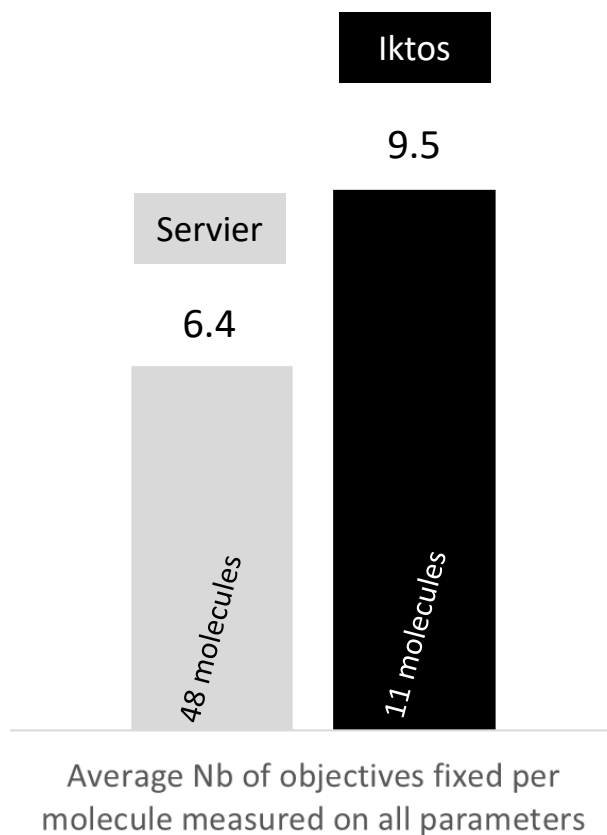
11 compounds were synthesized and tested

Final results af



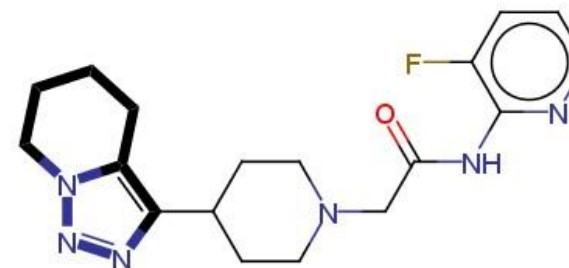
For the majority of objectives, Iktos molecules have a much higher "in blueprint" rate

MPO sum



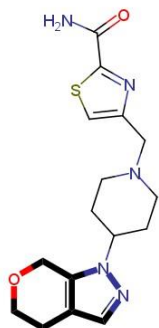
Best AI designed compound

Activity	5-HT2A	5-HT2B	a1	D1	Na _v 1.2	hERG	RLM	HLM	Caco-2 FAbs	Caco-2 Efflux
	83	7	18	7	-9	2	3	57	75	97



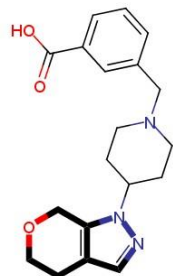
Presence of a [1,2,3]triazolo[1,5-a]pyridine moiety which appears only 6 times in the initial dataset, all having issues on CACO-2 (Permeability & Efflux)

AI designed

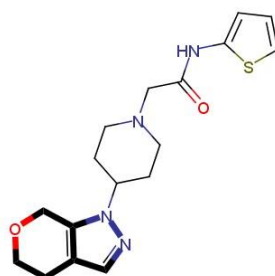


Active 9/11

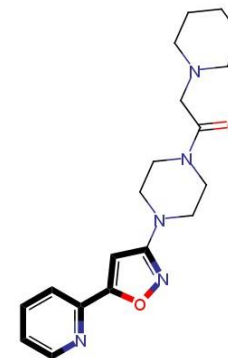
Presence of a pyrano[3,4-c]pyrazole moiety which appears 5 times in the initial dataset



Active 9/11



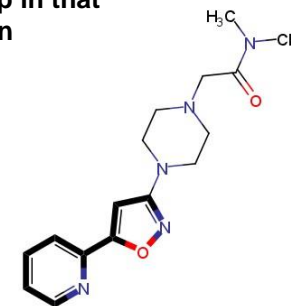
Active 10/11



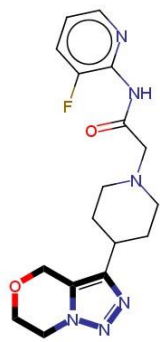
Borderline* 10/11

Presence of a 2-(1,2-oxazol-5-yl)pyridine moiety which appears 13 times in the initial dataset

First introduction of an aliphatic group in that position

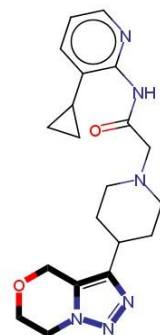


Inactive 8/11

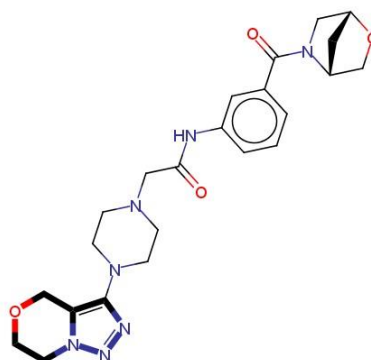


Borderline* 10/11

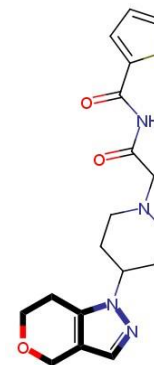
Presence of a [1,2,3]triazolo[4,3-c][1,4]oxazine moiety which appears 4 times in the initial dataset



Inactive 8/11

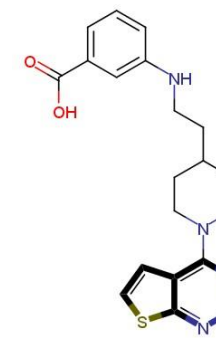


Active 9/11



Active 8/11

Presence of a pyrano[4,3-c]pyrazole moiety which appears 5 times in the initial dataset

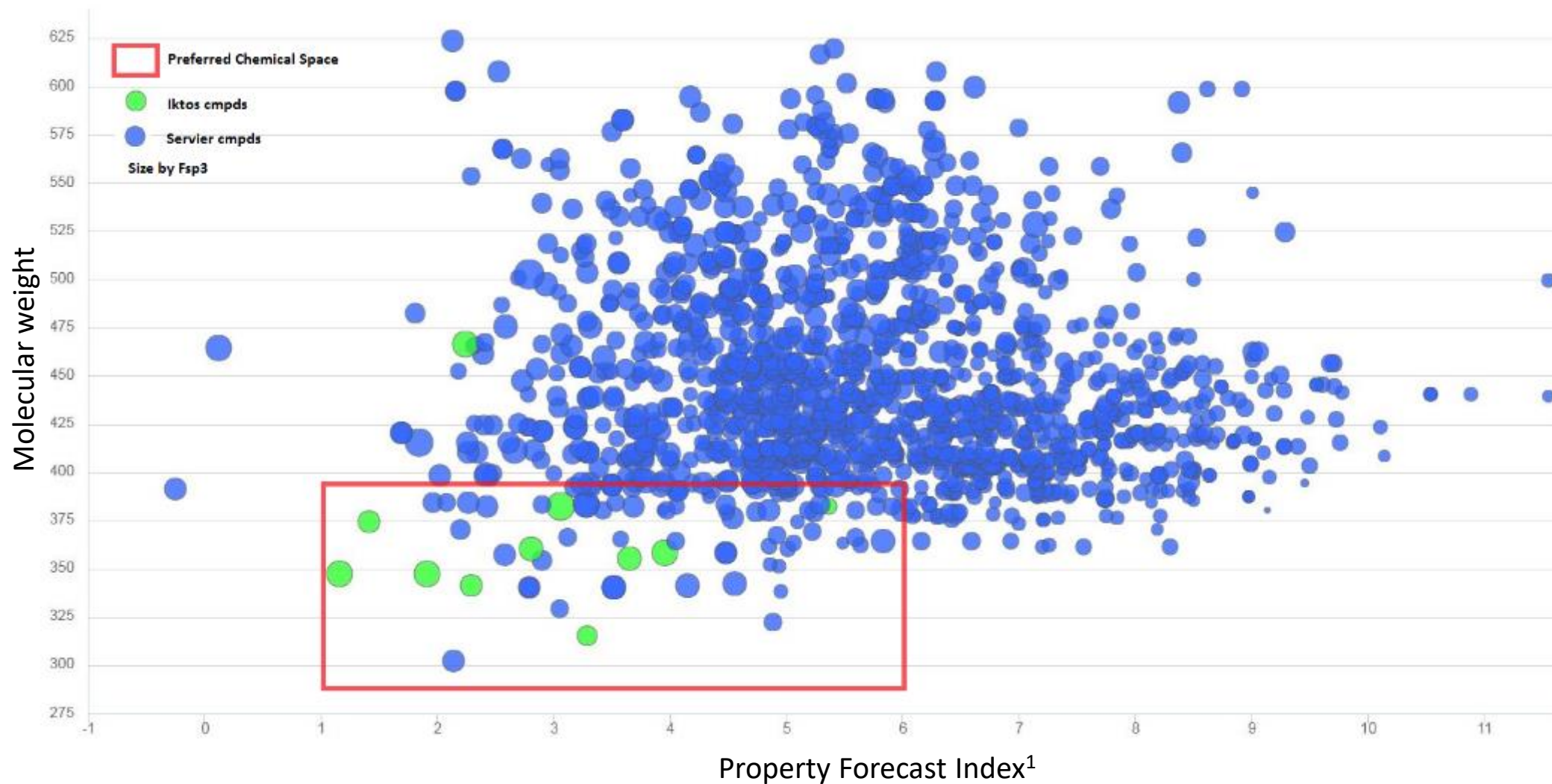


Inactive 9/11

Presence of a thieno[2,3-d]pyrimidine moiety which was absent from the initial dataset

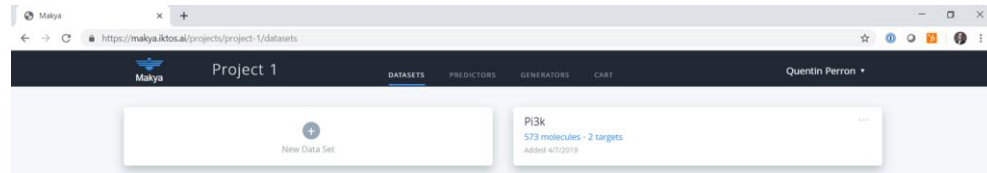
* In the margin error of the Activity assay

Drug-lik

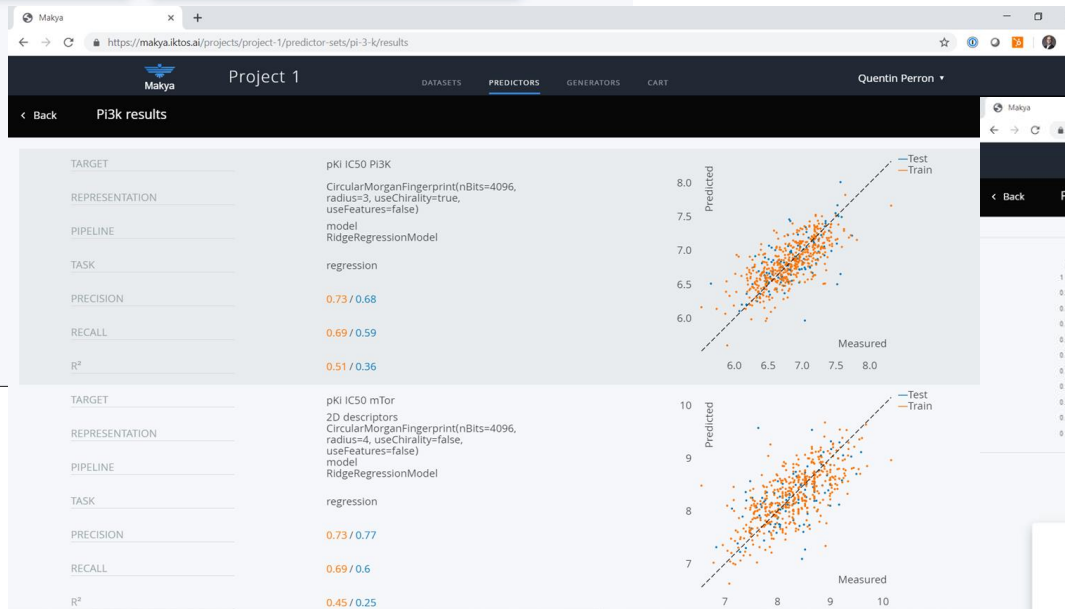


1 - Paul D. Leeson and Robert J. Young Molecular Property Design: Does Everyone Get It? *ACS Med Chem Lett.* **2015**, 6(7), 722–725.

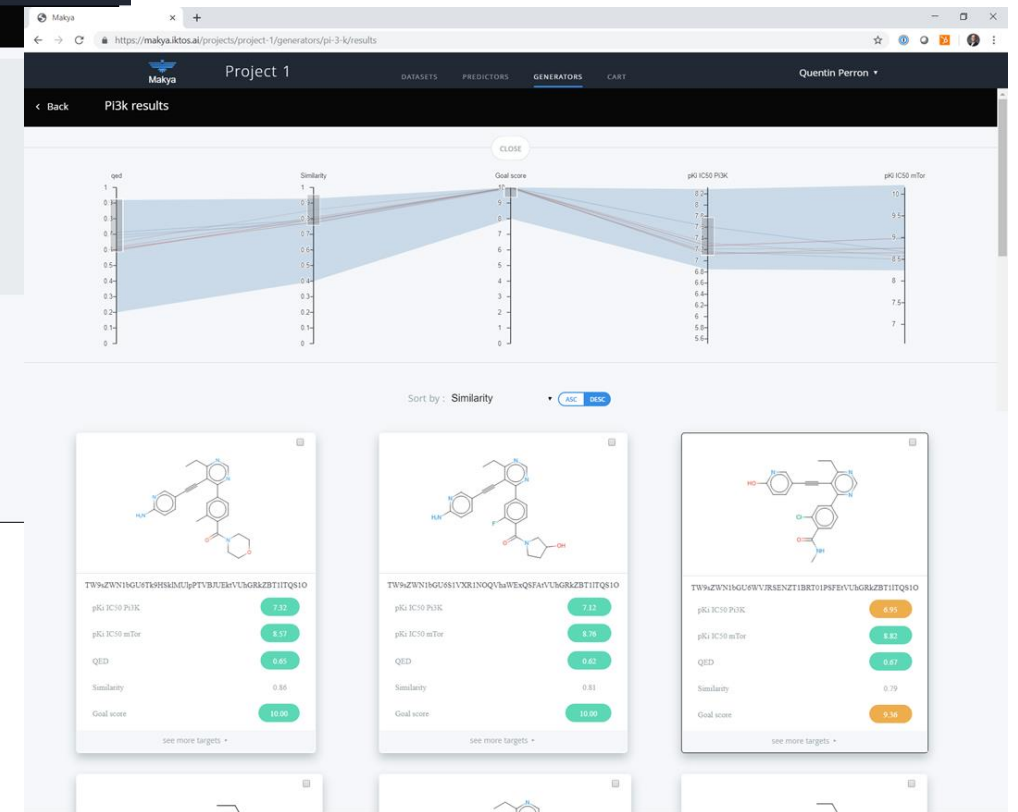
Generative mo



Dataset upload



AutoML module



“ideal” in silico propositions

Conclu

- ⊠ Optimal *in silico* propositions in only a few hours
- ⊠ An approach which is complementary to the propositions made by the medicinal chemist
- ⊠ High potential of the technology to accelerate early stage lead optimization – considering several compound design, synthesis and test iterations
- ⊠ Only local models were used (using the project's data) – likely benefit of integrating in house global predictive models into the process
- ⊠ Future developments: coupling generative models with structure-based modeling

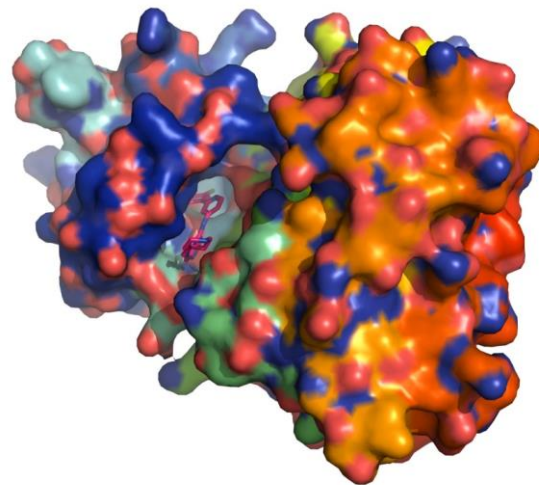


Perspectives

Design with a score

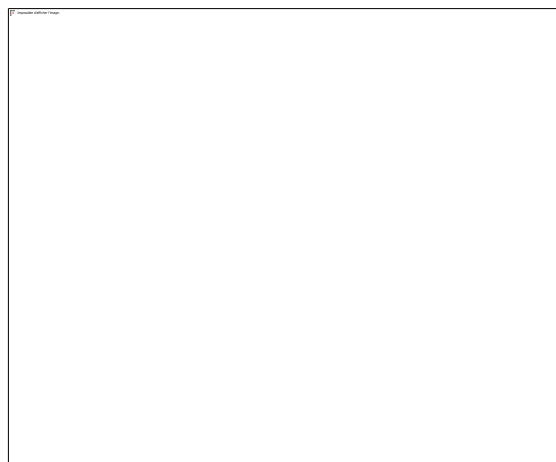


Generator

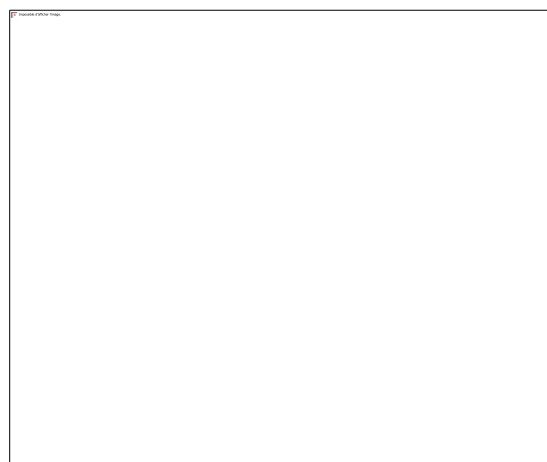


Docking

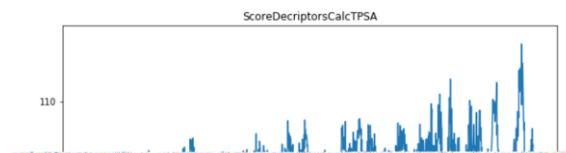
- ✓ High predicted value
- ✓ Important interactions are present
- ✓ All important PhysChem properties are present



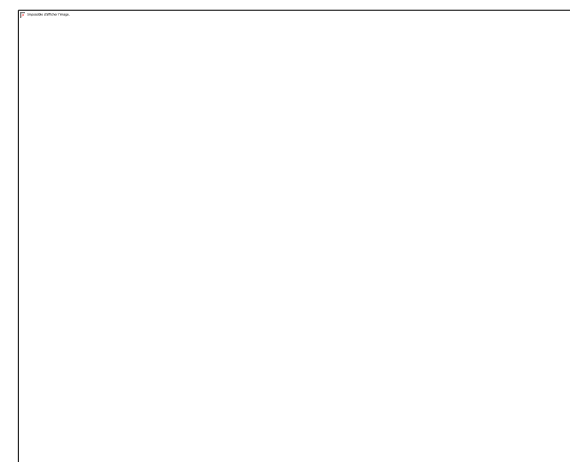
Docking Score



Contact Score

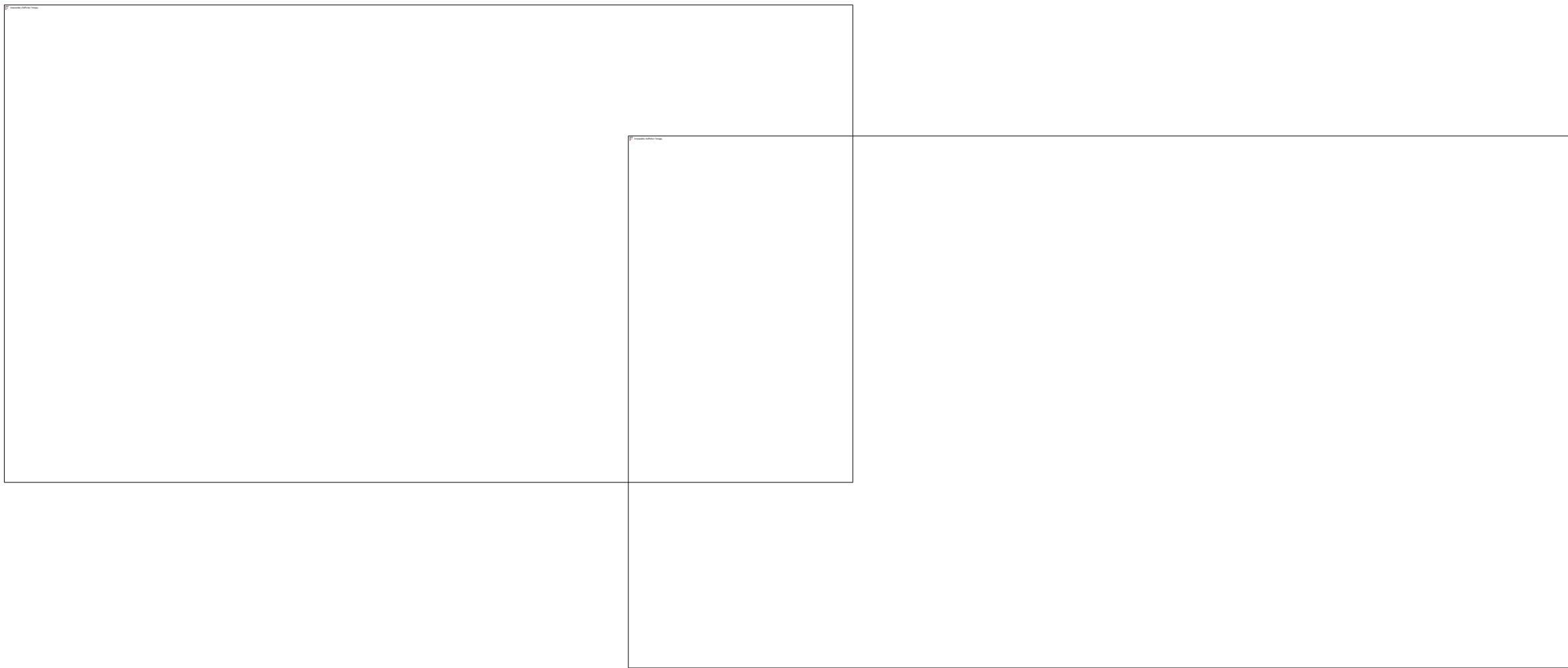


TPSA



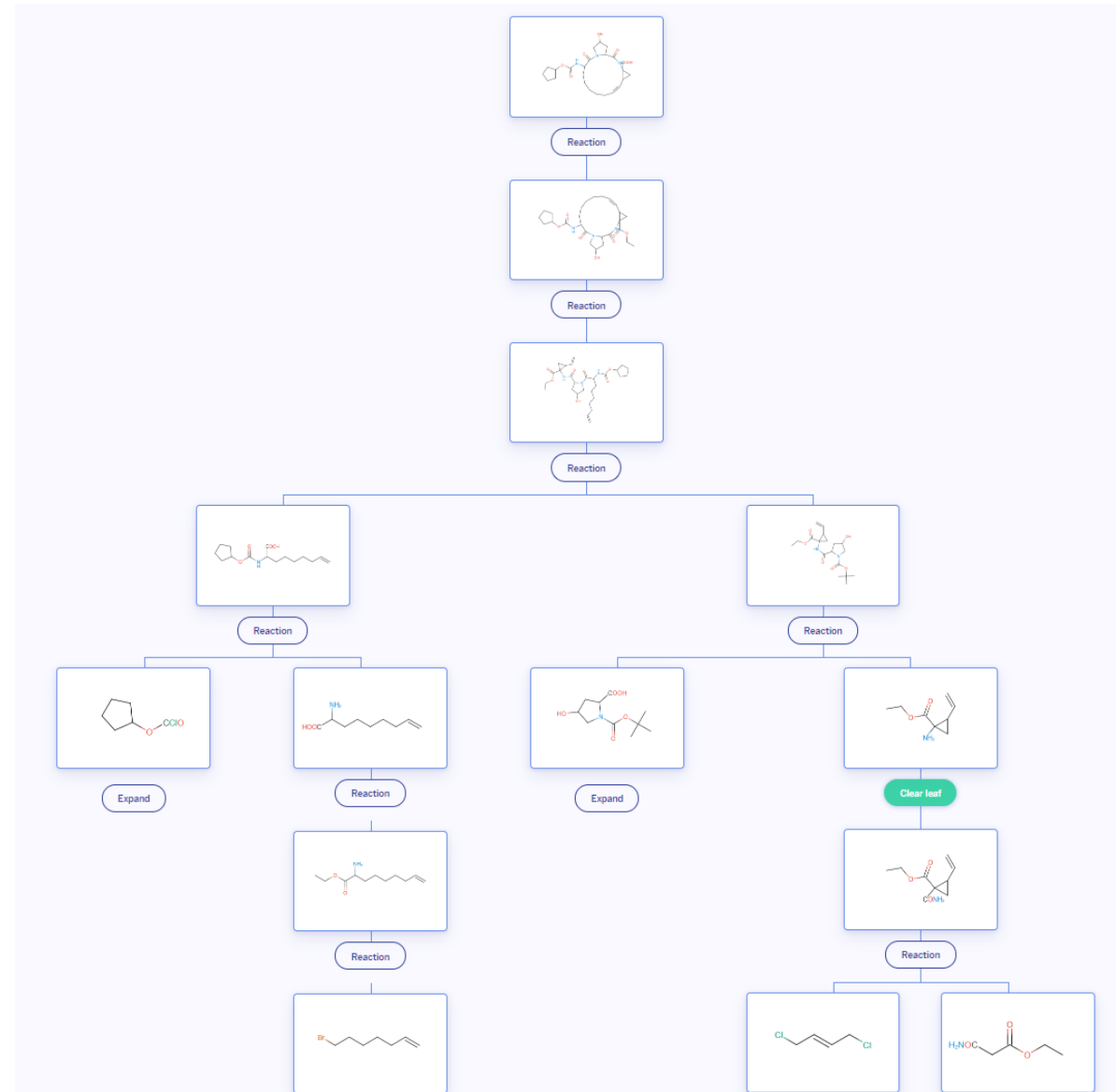
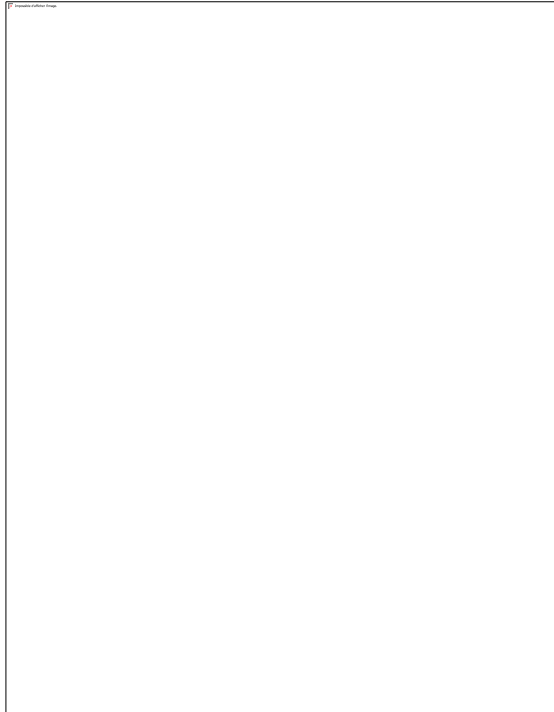
LogP

retrosynthesis



Freely accessible Beta version coming soon!

Ciluprevir int



Acknowled



The Iktos team!



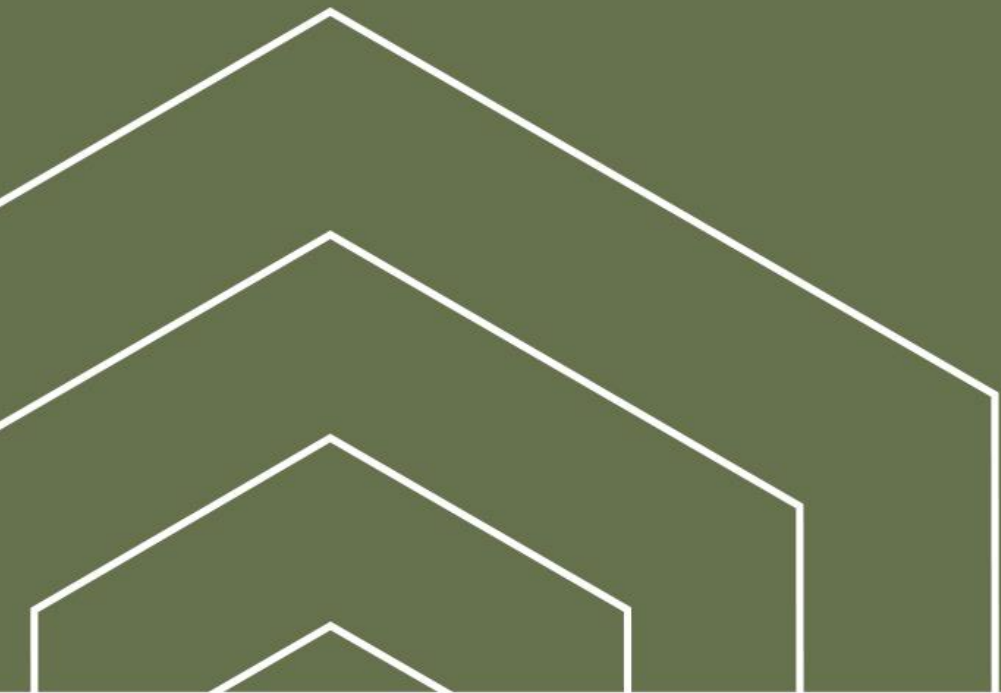
Olivier Mirguet, Anne Rojas, Pierre Ducrot, Arnaud Gohier, Marie-Pierre Bourguignon, Patricia Sansilvestri, Philippe Gloanec, Françoise Gellibert



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Back up slides



IKTOS